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<!--StartFragment-->RESULT 5
HUMORF08
LOCUS      HUMORF08              3694 bp      mRNA      linear      PRI 24-JAN-2003
DEFINITION Homo sapiens KIAA0022 mRNA, complete cds.
ACCESSION  D14664
VERSION    D14664.1   GI:285952
KEYWORDS   .
SOURCE     Homo sapiens (human)
  ORGANISM Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Euarchontoglires; Primates; Haplorrhini;
            Catarrhini; Hominidae; Homo.
REFERENCE  1
  AUTHORS  Nomura,N., Miyajima,N., Sazuka,T., Tanaka,A., Kawarabayasi,Y.,
            Sato,S., Nagase,T., Seki,N., Ishikawa,K. and Tabata,S.
  TITLE    Prediction of the coding sequences of unidentified human genes. I.
            The coding sequences of 40 new genes (KIAA0001-KIAA0040) deduced by
            analysis of randomly sampled cDNA clones from human immature
            myeloid cell line KG-1
  JOURNAL  DNA Res. 1 (1), 27-35 (1994)
  PUBMED   7584026
REFERENCE  2
  AUTHORS  Nomura,N., Miyajima,N., Sazuka,T., Tanaka,A., Kawarabayasi,Y.,
            Sato,S., Nagase,T., Seki,N., Ishikawa,K. and Tabata,S.
  TITLE    Prediction of the coding sequences of unidentified human genes. I.
            The coding sequences of 40 new genes (KIAA0001-KIAA0040) deduced by
            analysis of randomly sampled cDNA clones from human immature
            myeloid cell line KG-1 (supplement)
  JOURNAL  DNA Res. 1 (1), 47-56 (1994)
  PUBMED   7584028
REFERENCE  3 (bases 1 to 3694)
  AUTHORS  Ohara,O., Nagase,T., Kikuno,R. and Nomura,N.
  TITLE    Direct Submission
  JOURNAL  Submitted (11-NOV-1992) Osamu Ohara, Kazusa DNA Research Institute;
            1532-3, Yana, Kisarazu, Chiba 292-0812, Japan
            (E-mail:cdnainfo@kazusa.or.jp, Tel:+81-438-52-3913)
FEATURES             Location/Qualifiers
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                       /cell_type="Myeloblast"
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ORIGIN

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Query Match 98.8%; Score 3694; DB 5; Length 3694;
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Qy	107	CGCGGACTGTCCTTCATCTACTTGGATTTCAGTTCCAAGACAGTTGTTACATTTTTCTCCA	166
Db	61	CGCGGACTGTCCTTCATCTACTTGGATTTCAGTTCCAAGACAGTTGTTACATTTTTCTCCA	120
Qy	167	AGAAGCCATCAAAGTAGAAAGCATAGAGGATGTCAGAAATCAGTGTACTGACCATGGAGC	226
Db	121	AGAAGCCATCAAAGTAGAAAGCATAGAGGATGTCAGAAATCAGTGTACTGACCATGGAGC	180
Qy	227	GGACATGATAAGCATAACATAATGAAGAAGAAAATGCTTTTATACTGGATACTTTGAAAAA	286
Db	181	GGACATGATAAGCATAACATAATGAAGAAGAAAATGCTTTTATACTGGATACTTTGAAAAA	240
Qy	287	GCAATGGAAAGGCCCAGATGATATCCTACTAGGCATGTTTTATGACACAGATGATGCGAG	346
Db	241	GCAATGGAAAGGCCCAGATGATATCCTACTAGGCATGTTTTATGACACAGATGATGCGAG	300
Qy	347	TTTCAAGTGGTTTGATAATTCAAATATGACATTTGATAAGTGGACAGACCAAGATGATGA	406
Db	301	TTTCAAGTGGTTTGATAATTCAAATATGACATTTGATAAGTGGACAGACCAAGATGATGA	360
Qy	407	TGAGGATTTAGTTGACACCTGTGCTTTTCTGCACATCAAGACAGGTGAATGGAAAAAAGG	466
Db	361	TGAGGATTTAGTTGACACCTGTGCTTTTCTGCACATCAAGACAGGTGAATGGAAAAAAGG	420
Qy	467	AAATTGTGAAGTTTCTTCTGTGGAAGGAACACTATGCAAAACAGCTATCCCATACAAAAG	526
Db	421	AAATTGTGAAGTTTCTTCTGTGGAAGGAACACTATGCAAAACAGCTATCCCATACAAAAG	480
Qy	527	GAAATATTTATCAGATAACCACATTTTAATATCAGCATTGGTGATTGCTAGCACGGTAAT	586
Db	481	GAAATATTTATCAGATAACCACATTTTAATATCAGCATTGGTGATTGCTAGCACGGTAAT	540
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Db	601	CACCACAGTTTTTTCAACCGCACCCCAATCACCTTATAATGAAGACTGTGTTTTGGTAGT	660
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Qy	827	TGGATATTAGAGCTTTAATGGTATTCCTTATTCCAGTAACATTTTTATGTACTCATCTGC	886
Db	781	TGGATATTAGAGCTTTAATGGTATTCCTTATTCCAGTAACATTTTTATGTACTCATCTGC	840
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Db	841	 TGTGAAAAGTCTTTAGGTTTCATTAAAAAAACAGGTTTTAGAAATGATCTTAGATCTAATA	900
Qy	947	 TAGTGATTTTAAGCATCCCGTCAAAGGCAGAACTGTCACTTGAATGAAGGAAAGCTTAA	1006
Db	901	 TAGTGATTTTAAGCATCCCGTCAAAGGCAGAACTGTCACTTGAATGAAGGAAAGCTTAA	960
Qy	1007	 AGCCCAAGCAGATAAAAAATAAAAGCCCAGCCTATTTGTCTTGCCTGCTGTATCTTCCCTA	1066
Db	961	 AGCCCAAGCAGATAAAAAATAAAAGCCCAGCCTATTTGTCTTGCCTGCTGTATCTTCCCTA	1020
Qy	1067	 TTTAGTTGACCCACTTTAGTTTATATGTTTATTAGTAAACATGAAATGGGGAATAAGTGA	1126
Db	1021	 TTTAGTTGACCCACTTTAGTTTATATGTTTATTAGTAAACATGAAATGGGGAATAAGTGA	1080
Qy	1127	 TTTTAAGTACATCCCATACATTTAAATATCTTTGATAATTGTTATTTTTTTGGCAGATAA	1186
Db	1081	 TTTTAAGTACATCCCATACATTTAAATATCTTTGATAATTGTTATTTTTTTGGCAGATAA	1140
Qy	1187	 TTCCTCTAGAATGTGTATCTTTTTATGATTTAGATGAAGAAAATTTTACAACCTTTTAACA	1246
Db	1141	 TTCCTCTAGAATGTGTATCTTTTTATGATTTAGATGAAGAAAATTTTACAACCTTTTAACA	1200
Qy	1247	 CCCCACACCAATTTTAGTTTCATTACTTTTACACACACCATTTTATCACAAATGACTCAA	1306
Db	1201	 CCCCACACCAATTTTAGTTTCATTACTTTTACACACACCATTTTATCACAAATGACTCAA	1260
Qy	1307	 GTTTTAATGAATGTTTATAAATTATTTGAAACAAAATATGATCGCTGTGTCCAGGATGGC	1366
Db	1261	 GTTTTAATGAATGTTTATAAATTATTTGAAACAAAATATGATCGCTGTGTCCAGGATGGC	1320
Qy	1367	 ATAGAGAAAGCTGGCAATTAGGTTAACACTTACATATTATAGTGCCCTTTAAGGATTTT	1426
Db	1321	 ATAGAGAAAGCTGGCAATTAGGTTAACACTTACATATTATAGTGCCCTTTAAGGATTTT	1380
Qy	1427	 TCTCTTGCCACCATAACCTTTTGTACTTTCCCTTATACAAGATGTATCTCATTCTCCTCAA	1486
Db	1381	 TCTCTTGCCACCATAACCTTTTGTACTTTCCCTTATACAAGATGTATCTCATTCTCCTCAA	1440
Qy	1487	 GCATTTATAAATTTTTCTTCAATGACATGAAAACGTGTGCAAGCAAAAACCGAAGAAAAA	1546
Db	1441	 GCATTTATAAATTTTTCTTCAATGACATGAAAACGTGTGCAAGCAAAAACCGAAGAAAAA	1500
Qy	1547	 CACTTAAGTACAACCTGTAGTGACAGTGATCAAAGTTTTTCAGTGCATTTATTGTACATTTT	1606
Db	1501	 CACTTAAGTACAACCTGTAGTGACAGTGATCAAAGTTTTTCAGTGCATTTATTGTACATTTT	1560
Qy	1607	 AAGAAAAAGGTGAAAATCATTTGGGGAGTAAAAAAATGAAAAAGCTGAAACGAGTAATTT	1666
Db	1561	 AAGAAAAAGGTGAAAATCATTTGGGGAGTAAAAAAATGAAAAAGCTGAAACGAGTAATTT	1620
Qy	1667	 TCCTCACCATCAATAAACCAAAAAACAGGAAAGATAAAGAATGTATAAATTTACGTAAT	1726
Db	1621	 TCCTCACCATCAATAAACCAAAAAACAGGAAAGATAAAGAATGTATAAATTTACGTAAT	1680
Qy	1727	 TAGTCACGTATCACTTATCAATGGGGATACGTTCTAAGAAATGCATAGTTAGGGAATCTT	1786
Db	1681	 TAGTCACGTATCACTTATCAATGGGGATACGTTCTAAGAAATGCATAGTTAGGGAATCTT	1740
Qy	1787	 GTGTGAAAATCAGCTTGTATTTACACAAACCCAGATGGTAGAGCCTATTTTGTCCCAAAC	1846

Db	1741	GTGTGAAAATCAGCTTGTATTTACACAAACCCAGATGGTAGAGCCTATTTTGTCCCAAAC	1800
Qy	1847	CTACACAGCATGTTACTGTGCTGAATACTGCAGACAATTGTAACACAATATTTGTGTATC	1906
Db	1801	CTACACAGCATGTTACTGTGCTGAATACTGCAGACAATTGTAACACAATATTTGTGTATC	1860
Qy	1907	TAAATATAGAAAAGGTACAGTAAAAATATGGTCTACTAAGGAAACACTGTTCTATATGTG	1966
Db	1861	TAAATATAGAAAAGGTACAGTAAAAATATGGTCTACTAAGGAAACACTGTTCTATATGTG	1920
Qy	1967	GTCCATTACTGACTGAAGTATACTGTCTAGAAGTCTGAGGCTCAAAGAAAAGTAATCCCT	2026
Db	1921	GTCCATTACTGACTGAAGTATACTGTCTAGAAGTCTGAGGCTCAAAGAAAAGTAATCCCT	1980
Qy	2027	CTTCTGAATCCACACCCCATCAATTATCTTACTTTCTTCTGCGGAGATAGATAGATATAC	2086
Db	1981	CTTCTGAATCCACACCCCATCAATTATCTTACTTTCTTCTGCGGAGATAGATAGATATAC	2040
Qy	2087	TATCTCACTAGCTTGACTAATGGCAACAAAGTTCCAGCTTGTGTAGTCTCTTTTTATTGA	2146
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Db	2221	ACACCAGCTGTTACACAAGCACAAGCATGCTCTGTAAGAGCTTTACATTTCTGAGATTTT	2280
Qy	2327	GTATAGTGATTGAGATGTCTATTTTATTATTGATAGACTATTACTAATGTCAATATTGAA	2386
Db	2281	GTATAGTGATTGAGATGTCTATTTTATTATTGATAGACTATTACTAATGTCAATATTGAA	2340
Qy	2387	CACTACCCTGGAATTCTGCCTGGTTTTCTACCCAAATTGTACCACTCCTTGAAGAACT	2446
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Qy	2447	ACAGGCACAGTAAAAAAATATGGCGTATTATGTGAACTAAAAGAGTTCTAAAGGAGTTCT	2506
Db	2401	ACAGGCACAGTAAAAAAATATGGCGTATTATGTGAACTAAAAGAGTTCTAAAGGAGTTCT	2460
Qy	2507	TAAAGGAGTGGTAGAATTTGGGTAGGAAAGTGATTAAGTCCAACCTAAAACCAACAGTCT	2566
Db	2461	TAAAGGAGTGGTAGAATTTGGGTAGGAAAGTGATTAAGTCCAACCTAAAACCAACAGTCT	2520
Qy	2567	CAAACGTCTACAACCTACAATGTCCAATGAGCCACTAGCCACATGAGGCTATTTAAGTAAA	2626
Db	2521	CAAACGTCTACAACCTACAATGTCCAATGAGCCACTAGCCACATGAGGCTATTTAAGTAAA	2580
Qy	2627	TTTAGTTTAAAATCCAGTTTTTCGAATTACATTAGCCACATTGTCAAGTGTTCAAATCACA	2686
Db	2581	TTTAGTTTAAAATCCAGTTTTTCGAATTACATTAGCCACATTGTCAAGTGTTCAAATCACA	2640
Qy	2687	GGTGGTTAGTGGCTACTGTACTGGGCAACATACATTATAGAACATTTTCATTATAGGAAG	2746
Db	2641	GGTGGTTAGTGGCTACTGTACTGGGCAACATACATTATAGAACATTTTCATTATAGGAAG	2700

Qy	2747	TTTTATTGGGCAGTGCTGCTCTTAAATCCTACCTTCCACTCAACTCCCATACAACTTTCT	2806
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Qy	2807	TTTGTACATTTTGATACTTTCTACCTAATGGCAGCTCTTCCAAAATAGCTGCTTTAAACT	2866
Db	2761	TTTGTACATTTTGATACTTTCTACCTAATGGCAGCTCTTCCAAAATAGCTGCTTTAAACT	2820
Qy	2867	CTGATTTAATTTTCAATATTTGGTTTCATTTTCAACAGGCCAAGAGGCCTCTGGTAATG	2926
Db	2821	CTGATTTAATTTTCAATATTTGGTTTCATTTTCAACAGGCCAAGAGGCCTCTGGTAATG	2880
Qy	2927	AAGTGCTATATATATATATATATGACGGAGTCTCACTGTGCTGCCCAGGCTACAGTGCAG	2986
Db	2881	AAGTGCTATATATATATATATATGACGGAGTCTCACTGTGCTGCCCAGGCTACAGTGCAG	2940
Qy	2987	TGGCTCGATCTTGGCTCTCTCCAATCTCCGCCTTGCAGGTTTTCAAGCAATTCTCCTGCC	3046
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Db	3001	TCAGCCTCCTTAGTAGCTGGGACCACAGACATCTGTCAACCACACCCAGCTAACTTTTTGT	3060
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Qy	3527	TCTACCGTCATTTACATGATAATCTGAAGCTAATATGACAATATTTAAATACTAAGTGGT	3586
Db	3481	TCTACCGTCATTTACATGATAATCTGAAGCTAATATGACAATATTTAAATACTAAGTGGT	3540
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